



Date: Friday, September 19th, 2014 Time: 12:00 PM - 1:00 PM EDT Presenter: Bryce Christensen, Ph.D. Statistical Geneticist, Golden Helix

In this webinar, you will learn about the latest bioinformatics analysis and tools for analyzing **GWAS and next-generation DNA sequencing data**, including practical applications for identifying and interpreting candidate functional variants. Dr. Bryce Christensen will present updated workflows with Golden Helix's newest products, SNP & Variation Suite 8 (SVS) and the revolutionary browser, GenomeBrowse.

Register at: www2.gotomeeting.com/ register/521827866





## WEBINAR: Advancing Sequencing and Genotype Analysis at the University of Florida

## Agenda:

- 1. Introduction to SVS and GenomeBrowse
  - Overview of the workspace: (spreadsheets, annotations, visualizations, etc.)
  - Working efficiently with big data

## 2. GWAS for Agrigenomics

- Data Management & Quality Assurance
- Analysis (SNP, CNV, Mixed Linear Model)
- Genomic Predictions (gBLUP)
- Visualization (QQ and Manhattan Plots)

## 3. DNA-Sequencing

- Exome filtering and variant sharing between samples
- Sample pattern filtering
- Variant classification
- Interrogating VCF and BAM files
- Visualizing data using GenomeBrowse

4. Q&A

